

SEQUENCE LISTING

<110> YANAI, Koji OKAKURA, Kaoru YASUDA, Shohei WATANABE, Manabu MIYAMOTO, Koichi MIDOH, Naoki MURAKAMI, Takeshi

- <120> Transformants producing secondary metabolites modified with functional groups, and novel biosynthesis genes
- <130> 2002-0451A/LC/00144
- <140> NEW
- <141> 2002-03-29
- <150> JP 11-276314
- <151> 1999-09-29
- <160> 23
- <170> PatentIn Ver. 2.1
- <210> 1
- <211> 2061
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 Phe Gln Tyr Ile Gly Glu Ala Thr Gly Gln Pro Pro Val Val Val Pro
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- gtc gtg tcc ccg ggc ccc ggc agc ccc gac cgg gaa cgg gac ttc gga 192 Val Val Ser Pro Gly Pro Gly Ser Pro Asp Arg Glu Arg Asp Phe Gly
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							gcc Ala									672
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Cys Leu Gly His Gln Gly Ile Ala Gln Leu Phe Gly Gly Thr Val Gly
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90
95

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Tyr His Ser Leu Ala Ala Thr Asp Leu Pro Asp Glu Leu Glu Pro Leu 130 135 140

Ala Trp Ser Asp Asp Gly Val Val Met Gly Leu Arg His Arg Glu Lys 145 150 155 160

Pro Leu Trp Gly Val Gln Phe His Pro Glu Ser Ile Gly Ser Asp Phe 165 170 175

Gly Arg Glu Ile Met Ala Asn Phe Arg Asp Leu Ala Leu Ala His His 180 185 190

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Arg Val Asp Val Leu Pro Asp Ala Glu Glu Val Arg Arg Gly Cys Leu 210 215 220

Pro Gly Glu Gly Thr Thr Phe Trp Leu Asp Ser Ser Ser Val Leu Glu 225 235 240

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Glu Tyr Leu Thr Tyr Arg Val Ala Asp Gly Val Val Ser Val Arg Gly 260 265 270

Ser Asp Gly Thr Thr Thr Arg Thr Arg Arg Pro Phe Phe Asn Tyr Leu 275 280 285

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Phe Glu Phe Asn Leu Gly Tyr Val Gly Tyr Leu Gly Tyr Glu Leu Lys 305 310 315

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Ala Phe Leu Phe Ala Asp Arg Ala Ile Ala Leu Asp His Gln Glu Gly 340 345 350

Cys Cys Tyr Leu Leu Ala Leu Asp Arg Gly His Asp Asp Gly Ala 355 360 365

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Pro Gly Arg Val Ser Leu Val Lys Asp Arg Ala Ala Arg Tyr Ala Ala

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ggc gtc a Gly Val T 145		Leu I												480
gta cgg c Val Arg I														528
gcc ctg a Ala Leu T														576
ctc ggc g Leu Gly V 1														624
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gtg tac g Val Tyr G 225		Ile (720
cgg gcg c Arg Ala I														768
gac ccg g Asp Pro A														816
ccc ggg g Pro Gly G 2														864
ctc cgc c Leu Arg A 290	egg ctc Arg Leu	atg g Met (Gly	ccg Pro 295	gag Glu	ctc Leu	gcg Ala	gcg Ala	ggc Gly 300	cag Gln	gac Asp	cac His	tgc Cys	912
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295

320 305 310 315 Asp Arg <210> 7 <211> 32 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: PCR primer for the pabAB gene <400> 7 32 ggggggatcc tatgcgcacg cttctgatcg ac <210> 8 <211> 32 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: PCR primer for the pabAB gene ggggggatcc tcatcgggcg cccgccactg cg 32 <210> 9 <211> 30 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: PCR primer for the papA gene <400> 9 30 ggtgatcata tgcgcacgct tctgatcgac <210> 10 <211> 30 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: PCR primer for

Gln Glu Leu Phe Arg Thr Leu His Arg Thr Asp Asp Glu Gly Glu Lys

the papA gene

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